## SEQUENCE LISTING

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Ala Arg 65	Ala 50	35 Gly Leu	Leu Leu Leu	Ala Leu Ala	Val Ala 70	Asp 55 Gly	40 Met Pro	Val Ile Pro	Arg Ala Ser	Gln Thr 75	Lys 60 Gly	45 Lys Lys	Ala Met Thr	Ala Ala Cys	Gly Leu 80	
Ala Arg 65 Ala	Ala 50 Ala	35 Gly Leu Gly	Leu Leu Leu Ile Ser	Ala Leu Ala 85	Val Ala 70 Gln	Asp 55 Gly Glu	40 Met Pro Leu	Val Ile Pro Gly Ser	Arg Ala Ser 90	Gln Thr 75 Lys	Lys 60 Gly Val	45 Lys Lys Pro	Ala Met Thr Phe Thr	Ala Ala Cys 95	Gly Leu 80 Pro	
Ala Arg 65 Ala Met	Ala 50 Ala Leu	35 Gly Leu Gly	Leu Leu Leu Ile Ser 100	Ala Leu Ala 85 Glu	Val Ala 70 Gln Val	Asp 55 Gly Glu Tyr	40 Met Pro Leu Ser	Val Ile Pro Gly Ser 105	Arg Ala Ser 90 Glu	Gln Thr 75 Lys Val	Lys 60 Gly Val	45 Lys Lys Pro	Ala Met Thr Phe Thr 110	Ala Ala Cys 95 Glu	Gly Leu 80 Pro	

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140
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Glu Ser Thr Thr Gly Gly Tyr Ala Lys Ser Ile Ser His Val Ile Ile
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Gly Leu Lys Thr Val Lys Gly Thr Lys Gln Leu Lys Leu Asp Pro Ser
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Ile Tyr Asp Ala Leu Ile Lys Glu Lys Val Ala Val Gly Asp Val Ile
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            180
Tyr Ile Glu Ala Asn Ser Gly Ala Val Lys Arg Val Gly Arg Cys Asp
                            200
                                                205
Ser Phe Ala Thr Glu Tyr Asp Leu Glu Ala Glu Glu Tyr Val Pro Ile
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Pro Lys Gly Glu Val His Lys Lys Glu Ile Val Gln Asp Val Thr
                                        235
Leu His Asp Leu Asp Ala Ala Asn Ala Gln Pro Gln Gly Gly Gln Asp
                245
                                    250
Ile Leu Ser Leu Met Gly Gln Met Met Lys Pro Arg Lys Thr Glu Ile
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Thr Glu Lys Leu Arg Gln Glu Ile Asn Lys Val Val Asn Arg Tyr Ile
                            280
Asp Glu Gly Ile Ala Glu Leu Val Pro Gly Val Leu Phe Ile Asp Glu
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                                            300
Val His Met Leu Asp Ile Glu Cys Phe Ser Tyr Leu Asn Arg Ala Leu
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                                        315
Glu Ser Pro Leu Ser Pro Ile Val Ile Leu Ala Thr Asn Arg Gly Ile
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Cys Asn Val Arg Gly Thr Asp Met Thr Ser Pro His Gly Ile Pro Val
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Asp Leu Leu Asp Arg Leu Val Ile Ile Arg Thr Glu Thr Tyr Gly Pro
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Thr Glu Met Ile Gln Ile Leu Ala Ile Arg Ala Gln Val Glu Glu Ile
                        375
                                            380
Asp Ile Asp Glu Glu Ser Leu Ala Tyr Leu Gly Glu Ile Gly Gln Gln
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                                        395
Thr Ser Leu Arg His Ala Ile Gln Leu Leu Ser Pro Ala Ser Val Val
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Ala Lys Thr Asn Gly Arg Glu Lys Met Cys Lys Ala Asp Leu Glu Glu
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Val Ser Gly Leu Tyr Leu Asp Ala Lys Ser Ser Ala Arg Leu Leu Gln
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   Met Arg Ile Glu Glu Val Gln Ser Thr Ser Lys Lys Gln Arg Ile
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			atc Ile										156
	 _		gcg Ala				_	_	_	_	-		204
			gcg Ala										252
	_	_	gag Glu		 -	_	_			_		_	300
			tac Tyr 85										348
			aga Arg										396
	-		gag Glu	_	-				_		_		444
			tat Tyr										492
			ggg Gly										540
			aag Lys 165										588
			gga Gly										636
			gat Asp										684
			aag Lys										732
			gca Ala										780

ttg tcc Leu Ser 240		_		_	_	_			_	_		-			828
gaa aaa Glu Lys															876
gaa gga Glu Gly															924
cac ato His Met	_	_		-	_						-	-	_		972
agc cca Ser Pro	Leu														1020
aat gta Asn Val 320															1068
ctt cta Leu Leu															1116
gag ato Glu Met															1164
atg gat Met Asp															1212
tct tto Ser Let 385	Arg					_				_	_		-		1260
aag act Lys Thr 400															1308
agt ggg Ser Gly															1356
caa caa Gln Glr	_	_				taga	attto	ggg t	caco	ctgto	eg to	ggaa	gtct	е	1407
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gtcaccggtg caggaattgc cgtgtgtgtt ttttatcttg ctcatcggtg tccggaatct

1647

1707

1767

1827

1869

Met Ile Gln Ile Leu Ala Ile Arg Ala Gln Val Glu Glu Ile Asp Met 355 360 Asp Glu Glu Ser Leu Ala Tyr Leu Gly Glu Ile Gly Gln Gln Thr Ser 375 380 Leu Arg His Ala Ile Gln Leu Ile Ser Pro Ala Ser Val Val Ser Lys 390 395 Thr Asn Gly Arg Glu Lys Ile Cys Lys Ala Asp Leu Glu Glu Val Ser 405 410 Gly Leu Tyr Leu Asp Ala Lys Ser Ser Ala Arg Leu Leu Gln Glu Gln 420 425 Gln Glu Arg Tyr Ile Thr 435 <210> 11 <211> 36 <212> DNA

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 remove clones which have a poly(A) tail but no
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36